

The intron sequences between exons 2 - 3 and exons 18 - 19 are missing (introns: small letters, exons: capital letters). Small letters in the first exon indicate nucleotides that have not been unambiguously determined.

*Exon 1*

```

1  CGGGTGAATC CCGGCGCCGC GCCCGGACC CGCAGCTCCC TGCCTCCTC
51  CCTCCCAGCC GCTTTAACAC CCACACCCCA CAGTCTCTCC CACGsCCGCG
101 CCTTGGCGGC CCCACTGAAT CCCTACGCGG GGCCCAGCGG TACCGGGAGA
151 CCGGGCTAGC CTATGGGAGC GCCCAGATAA CGCGGGTTGG GGGCGCCCGC
201 GCCCCcATCC CCGCCAGCAAT GACTCGATCG CCGCCCCCTCA GAGAGCTGCC
251 CCCGAGTTAC ACACCCCCAG CTCGAACCGC AGCACCCAG gtgagtagag
301 ggggagctgg aagaaggaag agagcggagc caggtctgtc actcgggcct
351 ctgcaagggt tgtgatgtct tgaagtgccg agtgtcatta gatgtctgaa
401 ggcaagttag agccagcacc gcaagcaagt tgtgctgtgt tgtcgggtgtg
451 tctgtgccgg tgtctcctca tcgtctggcc agtgagaatg aatgtctgtg
501 ggttcacctc tgtgtccacc cgacgacagg tgtgtgtaca tatgtatcct
551 gctctcagaa aatgggccta tgccgccggg cgcggtgact cagcctgta
601 atcccaacac tgggaggtgt aggcaggcag attacctgag gtcaggagtt
651 cgagaccagc caggccaaca tggggaaact ctgtctctac taaaaataaa
701 aattagcagg gcgtggtggc gggcgctgt agtcccaact actcgggagg
751 ctgaggcagg agaattctct gaacctggga ggcggagggt gcagtcaagc
801 cgagatcaca ccaactgcact ccagccagg caacagagcg agatgcgtct
851 caaaaaaaaa aaaaaaaaaa aaaaggagag aaaacaaaaa gaaaagaaag
901 gaaaataggc ctatgccttc ctcagggtgt tgcctgggat ggtgggtgtt
951 acatottcca agtctgggcc tgtgtctgtg ttggtgctcc ctgtcccaca
1001 tccagaaatc aagaagcgag ggotgggcag cagatataca gggtgagaag

```

Fig. 1

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1051	ggaaggattt	catgcattgt	tacagtgatg	cctggctgac	ccttctcttt
1101	ccatcccaga	TCCTAGCTGG	GAGCCTGAAG	GCTCCACTCT	GGCTTCGTGC
1151	TTACTTCCAG	GGCCTGCTCT	TCTCTCTGGG	ATGCGGGATC	CAGAGACATT
1201	GTGGCAAAGT	GCTCTTTCTG	GGACTGTTGG	CCTTTGGGGC	CCTGGCATT
1251	GGTCTCCGCA	TGGCCATTAT	TGAGACAAAC	TTGGAACAGC	TCTGGGTAGA
1301	AGTGGGCAGC	CGGGTGAGCC	AGGAGCTGCA	TTACACCAAG	GAGAAGCTGG
1351	GGGAGGAGGC	TGCATACACC	TCTCAGATGC	TGATACAGAC	CGCACGCCAG
1401	GAGGGAGAGA	ACATCCTCAC	ACCCGAAGCA	CTTGGCCTCC	ACCTCCAGGC
1451	AGCCCTCACT	GCCAGTAAAG	TCCAAGTATC	ACTCTATGGG	AAG.....
1501	.....	.....	.....	.....	.....g
1551	tgagtctggc	tgagcccctg	agcagctggg	ggcgggcgt	gctgtggggg
1601	ttctggagtg	ggaatcccct	tcttctgctg	atctcctatg	cccctggcta
1651	ttgcagTCCT	GGGATTGAA	CAAAATCTGC	TACAAGTCAG	GAGTTCCCCT
1701	TATTGAAAAT	GGAATGATTG	AGCGGgtaag	tgtcctgaga	gggagtagag
1751	gcagaacttt	ttctgtagcg	tgggaggact	cagagaccga	gcaagcccca
1801	cagcctgcaa	tctgccccct	taaaactaag	gagggggatt	gcagagggca
1851	tcctacaaag	gttggtgggg	aggactgacg	tggcccgggg	tatccctggc
1901	agATGATTGA	GAAGCTGTTT	CCGTGCGTGA	TCCTCACCCC	CCTCGACTGC
1951	TTCTGGGAGG	GAGCCAAACT	CCAAGGGGGC	TCCGCCTACC	TGCCgtgagt
2001	gccactcctg	gggccctgct	tcctctcccg	ctggggactc	tcccagcaga
2051	aaggaggggt	ctgggggaatg	aggatgatca	aaaccttacc	aaggctcctaa
2101	ttacctccca	ggccaggaac	agagagcatg	ggcttcccca	aggctctctc
2151	cacatcctcc	ttctctttcc	ctctcaagga	aggaagacct	gacttattta
2201	cacaaaacta	aacacaaaga	tctgtaagat	ctgagcaaag	gagaaaaaga
2251	tccccacaaa	gaggctttgc	tgggggaaat	tacctaggtg	tttgctaagc
2301	cattgcccag	gccagaaaga	aaacctgcta	caggcatgtg	cctgctggtt
2351	gtatattaga	accaagcaca	cagcttggtg	aggaactcag	tggggccttt

Fig. 1 (cont.)

090707Z  
FM JCRC  
TO RUEADK/SECDEF  
INFO RUEADK/SECDEF

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2401 ctggggccctt tctatgtatt aggtaaccct gccctgatat t~~cg~~tctcagc  
 2451 cccttgtact cttctacagc t~~ca~~ctgtagc accctgggtg gcccatgcag  
 2501 cctggcagtt ctgagaagct gaggcttgca caccctccat atggaaggac  
 2551 aaatcggcag ataagaggag ggtggggtac agcatggcgc ccagcagca  
 2601 gtttggagcc tgggttttcg tccctgacct tcaccaacta taggcttttc  
 2651 cctcagCGGC CGCCCGGATA TCCAGTGGAC CAACCTGGAT CCAGAGCAGC  
 2701 TGCTGGAGGA GCTGGGTCCC TTTGCCTCCC TTGAGGGCTT CCGGGAGCTG  
 2751 CTAGACAAGG CACAGGTGGG CCAGGCCTAC GTGGGGCGGC CCTGTCTGCA  
 2801 CCCTGATGAC CTCCACTGCC CACCTAGTGC CCCCACCAT CACAGCAGGC  
 2851 AGgtgggttc caaccaggtc tgccagggaa aggctgtttt ccttcccttt  
 2901 cccttctctca tactcctgtg ttctggggga gctgactgct ctgtgccctg  
 2951 accccccaact tcctggccat tattaccctg ctcccacagt gccaggcccc  
 3001 caatgttcca ttcccattca gttatcctac ggagccctca agtgggtatat  
 3051 atgaatccct ttttcctttt ctaagcctag ataaggctgg acttcttttt  
 3101 tttttttttt ttgagtctca ctctgtcacc caggctggag tgcagtagtt  
 3151 cgatcttggc t~~ca~~ctgcaac ct~~cg~~ggtcaa gcaattctcc tgccttagcc  
 3201 tcctgagtag ctgggattac aggtgccac caccatgcc ggctaatttt  
 3251 tattagcctc ccaaagtgt gggattacag gcgtgagcca ctgcgcctgg  
 3301 ccaaggctgg actttttatc aaaatagact aatacagga aactaagaac  
 3351 acagcaggta agcatgaata tcatacctgg tttcccaggt ttctttgtgg  
 3401 ccctgcaaat gtggtacttt tttcagaatc cgccagttac accagctcct  
 3451 ccagaagcc tacttccagg cctctgcttc cccttggggc ttcctgtctg  
 3501 cgggatacta gctgttca~~ct~~ cctgcagagc agtcaagagg ct~~c~~agaatag  
 3551 ttacctacac tccagcccta ctgagcttca tggcagcgtg gttcctggag  
 3601 gtggaagccc agggacactc agttatccac ggccagggcc ttgagcatta  
 3651 acccctcctg ttccccctca GGGCTCCCAA TGTGGCTCAC GAGCTGAGTG  
 3701 GGGGCTGCCA TGGCTTCTCC CACAAATTCA TGC~~ACT~~GGCA GGAGGAATTG

Fig. 1 (cont.)

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3751 CTGCTGGGAG GCATGGCCAG AGACCCCCAA GGAGAGCTGC TGAGgtaggg  
 3801 tctcctctgg gagttggtga ggggactctg ttcattgagaa cccatactgt  
 3851 aatgccagge agctctggca aaaggccctt cacatccctc accaggtgtt  
 3901 tggggccagct ctgacccttg gttctccac acccccacca gGGCAGAGGC  
 3951 CCTGCAGAGC ACCTTCTTGC TGATGAGTCC CCGCCAGCTG TACGAGCATT  
 4001 TCCGGGGTGA CTATCAGACA CATGACATTG GCTGGAGTGA GGAGCAGGCC  
 4051 AGCACAGTGC TACAAGCCTG GCAGCGGCGC TTTGTGCAGg tgggtatgga  
 4101 caaggacaag ggggggtgcc tgaggccatt cctcctcct gccccctcct  
 4151 atccaccctg tttctccagc TGGCCCAGGA GGCCCTGCCT GAGAACGCTT  
 4201 CCCAGCAGAT CCATGCCTTC TCCTCCACCA CCCTGGATGA CATCCTGCAT  
 4251 GCGTTCTCTG AAGTCAGTGC TGCCCGTGTG GTGGGAGGCT ATCTGCTCAT  
 4301 Ggtgggtcctt gcacctggca cettgcccc accccacctc caaccagtgc  
 4351 ccaccctggg agcccttgag actgcccctt cccccacag CTGGCCTATG  
 4401 CCTGTGTGAC CATGCTGCGG TGGGACTGCG CCCAGTCCA GGGTTCCGTG  
 4451 GGCCCTGCCG GGGTACTGCT GGTGGCCCTG GCGGTGGCCT CAGGCCTTGG  
 4501 GCTCTGTGCC CTGCTCGGCA TCACCTTCAA TGCTGCCACT ACCCAGgtac  
 4551 gccaggactg cagggcagac tcagtgccag tcaccaggct tcacgggtcc  
 4601 tcagctgcc gctcctctgc cctccagGT GCTGCCCTTC TTGGCTCTGG  
 4651 GAATCGGCGT GGATGACGTA TTCCTGCTGG CGCATGCCTT CACAGAGGCT  
 4701 CTGCCTGGCA CCCCTCTCCA Ggtggggcct tgtccccag ggtcactctg  
 4751 aggcagctca gcttactggt taagagcctc ttggttcaag tgacccttgg  
 4801 gctgctaatag aacctcgggt cctctgtcc ccatctgtaa acaggggaaa  
 4851 taatagtgt gtgtcctaag ggttattgtt tggatcagtg aggtaactca  
 4901 agttgaatgc ttagaacagc ccatcatag tacatggtac ccaataaatg  
 4951 ctageccactg tgttatgact gccccacctc tgcaccccaa gttcctgagc  
 5001 ctcccccttca ctccactttg acacggcccc tcccttgtga cctgagggca  
 5051 ggtccccact ctgtcctggc agGAGCGCAT GGGCGAGTGT CTGCAGCGCA

Fig. 1 (cont.)

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5101 CGGGCACCAG TGTCGTACTC ACATCCATCA ACAACATGGC CGCCTTCCTC  
 5151 ATGGCTGCCC TCGTTCCCAT CCCTGCGCTG CGAGCCTTCT CCCTACAGGC  
 5201 GGCCATAGTG GTTGGCTGCA CCTTTGTAGC CGTGATGCTT GTCTTCCCAG  
 5251 CCATCCTCAG CCTGGACCTA CGGCGGCGCC ACTGCCAGCG CCTTGATGTG  
 5301 CTCTGCTGCT TCTCCAGgta ctgcggtgogc cccagccct tectcccgty  
 5351 acccacgcca gcctgtcccc tcaccagcat ttcaaggcac agacctgtca  
 5401 tccactctct acctcttcca gTCCCTGCTC TGCTCAGGTG ATTCAGATCC  
 EXON 13  
 5451 TGCCCCAGGA GCTGGGGGAC GGGACAGTAC CAGTGGGCAT TGCCCCACCTC  
 5501 ACTGCCACAG TTCAAGCCTT TACCCACTGT GAAGCCAGCA GCCAGCATGT  
 5551 GGTCACCATC CTGCCTCCCC AAGCCCACCT GGTGCCCCCA CCTTCTGACC  
 5601 CACTGGGCTC TGAGCTCTTC AGCCCTGGAG GGTCCACACG GGACCTTCTA  
 5651 GGCCAGGAGG AGGAGACAAG GCAGAAGGCA GCCTGCAAGT CCCTGCCCTG  
 5701 TGCCCGCTGG AATCTTGCCC ATTCGCCCCG CTATCAGTTT GCCCCGTTGC  
 5751 TGCTCCAGTC ACATGCTAAG gtaagactgg gcagagcagg gcagagactt  
 5801 agcatctctg ggcccagaag ggcagagagg gcttagtcca ctgcctgagg  
 5851 ggctgggggc agccctgggg tctccagctt agttgctaca tcccgcagGC  
 EXON 14  
 5901 CATCGTGCTG GTGCTCTTTG GTGCTCTTCT GGGCCTGAGC CTCTACGGAG  
 5951 CCACCTTGGT GCAAGACGGC CTGGCCCTGA CGGATGTGGT GCCTCGGGGC  
 6001 ACCAAGGAGC ATGCCTTCCT GAGCGCCCAG CTCAGGTACT TCTCCCTGTA  
 6051 CGAGGTGGCC CTGGTGACCC AGGGTGGCTT TGA CTACGCC CACTCCCAAC  
 6101 GCGCCCTCTT TGATCTGCAC CAGCGCTTCA GTTCCCTCAA GCGGGTGCTG  
 6151 CCCCCACCGG CCACCCAGGC ACCCCGCACC TGGCTGCACT ATTACCGCAA  
 6201 CTGGCTACAG Ggtgagaggg gaggagacgg gcagggaggg gtgctgcagg  
 6251 gagaaacgcc ctggggccac cagctaataa aaccctatcc tgggtctcccc  
 EXON 15  
 6301 cagGAATCCA GGCTGCCTTT GACCAGGACT GGGCTTCTGG GCGCATCACC  
 6351 CGCCACTCGA CCGCAATGGC TCTGAGGATG GGGCCCTGGC CTACAAGCTG  
 6401 CTCATCCAGA CTGGAGACGC CCAGGAGCTT CTGGATTTCa GCCAGgttgg

Fig. 1 (cont.)

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6451 gagagggctg gaggggtcca ctagtacagg ggctgcaggc ctcctgggccc  
 6501 caggccttca gccctctctg cctctgcagC TGACCACAAG GAAGCTGGTG  
 6551 GACAGAGAGG GACTGATTCC ACCCGAGCTC TTCTACATGG GGCTGACCGT  
 6601 GTGGGTGAGC AGTGACCCCC TGGGTCTGGC AGCCTCACAG GCCAACTTCT  
 6651 ACCCCCCACC TCCTGAATGG CTGCACGACA AATACGACAC CACGGGGGAG  
 6701 AACTTTTCGCA gtgagtcttg gggggagctc ggcaagagcc tcagcctcgc  
 6751 ccacacaagc cctgagcctg aggccctgcc cactctgccc cgtgctcacc  
 6801 gccctgtccc tctccctctt ctcccttccc ctccctccca cagTCCCGCC  
 6851 AGCTCAGCCC TTGGAGTTTG CCCAGTTCCC TTTCTGCTG CGTGGCCTCC  
 6901 AGAAGACTGC AGACTTTGTG GAGGCCATCG AGGGGGCCCG GGCAGCATGC  
 6951 GCAGAGGCCG GCCAGGCTGG GGTGCACGCC TACCCAGCG GCTCCCCCTT  
 7001 CCTCTTCTGG GAACAGTATC TGGGCCTGCG GCGCTGCTTC CTGCTGGCCG  
 7051 TCTGCATCCT GCTGGTGTGC ACTTTCCTCG TCTGTGCTCT GCTGCTCCTC  
 7101 AACCCTGGA CGGCTGGCCT CATAgtagt gcttgcagga gtggggacag  
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 7251 GGAACCTCTT GGTATCATGG GTTTCCTGGG CATCAAGCTG AGTGCCATCC  
 7301 CCGTGGTGAT CTTGTGGCC TCTGTAGGCA TTGGCGTTGA GTTCACAGTC  
 7351 CACGTGGCTC TGGGCTTCCT GACCACCCAG GGCAGCCGGA ACCTGCGGGC  
 7401 CGCCCATGCC CTTGAGCACA CATTGCCCC CGTGACCGAT GGGGCCATCT  
 7451 CCACATTGCT GGGTCTGCTC ATGCTTGCTG GTTCCCACTT TGACTTCATT  
 7501 GTAAG.....  
 7551 ..... gtagggaggg ctcggggcag ggaggcaggg ctcaggacag  
 7601 gcctgggctg actcccccca caccctaccc ctagGTACTT CTTTGC GGCG  
 7651 CTGACAGTGC TCACGCTCCT GGGCCTCCTC CATGGACTCG TGCTGCTGCC  
 7701 TGTGCTGCTG TCCATCCTGG GCCCGCCGCC AGAGgtgacc acapcctcgg  
 7751 caccatccct ctactcccag cccaaggagc ggggtaggga gaggcaaggg

Fig. 1 (cont.)

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7801 aagggacaga gccctgtggc ccacagacag gtacctcccc aacagggtgcc  
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 8151 GGAAAGCCCA GAGATCCTGA GTCCACCAGC TCCACAGGGA GCGGGGCTTA  
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 8301 tccagccct gatgagcccc cttgggtcccc tgcctgctact agctctggca  
 8351 acctcagttc caggggacca ggtccagcca ctgggtgaaa gagcagctga  
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 8551 agccaccgat ttgcacatcc aggcctgtgt gagcctgtat ctgtgtcact  
 8601 tgagagtga agetggcact tggggctgca gtgcagccct gtcccccttc  
 8651 ccaccccaca cactgcctg cccagctgac caagcctgag ggaccctcca  
 8701 gcacccttcc gtctgggtgac tcctgggcag gctctccata tccctgcccc  
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 8801 catacatgtt agctatgatg aaagttttat tttttaaaga atgaaatata  
 8851 ttctatgtga agctatgatg aaagttttat tttttaaaga atgaaatata  
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 9001 gtggggagtt tgcaagtga cattaactat tggaaggagc ttctctgggtg  
 9051 ccaggacctg aggtattagc ttctctagtt ctgggtggaa aagacccag  
 9101 attctggatt tttgtcatat acttggtaac atcatctgga ttaagtgcct

Fig. 1 (cont.)

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9151 actatacaaa acgataacaa attttgttgg tgtgaaatcc tactgggttc  
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 9251 gagccacat attccagcct gccgtctct ccagactcac ctccacctac  
 9301 ctgcttcacc cgcacgggaa acggcaaggc agaggggcaa agccatgcag  
 9351 caggtggaag gcgaggtgga ggcagatcag gaaagcagcc agttgaagca  
 9401 gagagaggtc aacaggggtct ggggagcttc tcaggagggtt tgtggacca  
 9451 gggaaaggag ccaggttcca gagcaacctc caaggcaaag gcctctgtaa  
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 9551 tcagttgctg gaactgctca gaaactgagg tgctagcagt tagtgaggac  
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 9901 aacaacaaaa aacgcctatt gcaattgaat ccacgctaaa atgcctaaaa  
 9951 agctcaagag aagcgggtag ttggcagaga acctagagta ggggggtgcaa  
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 10051 tcaccaagaa gggccttcta ggtggagcag agagagctca ccaggccaga  
 10101 atagtcaaaa gggggtcagc cctcagtgc acttaccagc ggagtaacc  
 10151 tgggcaagtt agccagcctc actaagcctc cccatcttca tctttccagG  
 XON 22  
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 10301 TTTCTCATAC GCCATTCCCT CTGCCTAGAA CACTTTCTCA CCTCCCCTTG  
 10351 ATGTGACCCC ATATCACCTT TCGAGGTGAA TTGGATCGGA TGCCATCTCC  
 10401 TCCAGGAGGG GTGGGGTCGT GCCTCCTGTG AGGTCCCAGT GCCCCTGAGT  
 10451 GTCTGTGCCC GTCTGTTTCC CCGTCCCTCT CTCTAAGCCC GGAGGCTTAC

Fig. 1 (cont.)

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10501 TCGGGGTAAG GACGGCGGGA CAGGACCTTA ACCGCTGGGA CGAACACCAG  
10551 CTCCGCAAAG GACTCCGCAC CCGGCGCCGC CCACGGGGTG CGGGTCCCAG  
10601 GAGGACCAGC AGAGAGGAGC ATAGGAGAGC AAAGGAGATC AGTGACCCAT  
10651 GGCTTCCCCG GTGGCGCGGA ACAGCCCGGA GCCGCCTGTG ATTTGCATAC  
10701 CCATGGTGCA CCACGAAAAG ATACCCTCAA GATGCTTGCA CTCCCTCTGT  
10751 GCGCGCATTT CTGCACTGTT TTAGAGCATG ATGCCTCTTA CACGCATCTG  
10801 TGTGCATAAA CTACATATAG GGAGTGCGTA CCACGCAGGC ATCCAACAAC  
10851 CATAAGTGTG TTAAGTGTTA GTTCTCCCTG CGAGGTTCGA AGCGGAAGTC  
10901 ACGAATATAC TCGGGTTTCT CTTCAAAGCG CATAAATCTT TCGCCTTTTA  
10951 CTAAAGATTT CCGTGGAGAG AAAGTTGTGA GTTTTTATTTC AATTTTTTTGA  
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11051 TACAAAAAAA AAAAAAAA

Fig. 1 (cont.)

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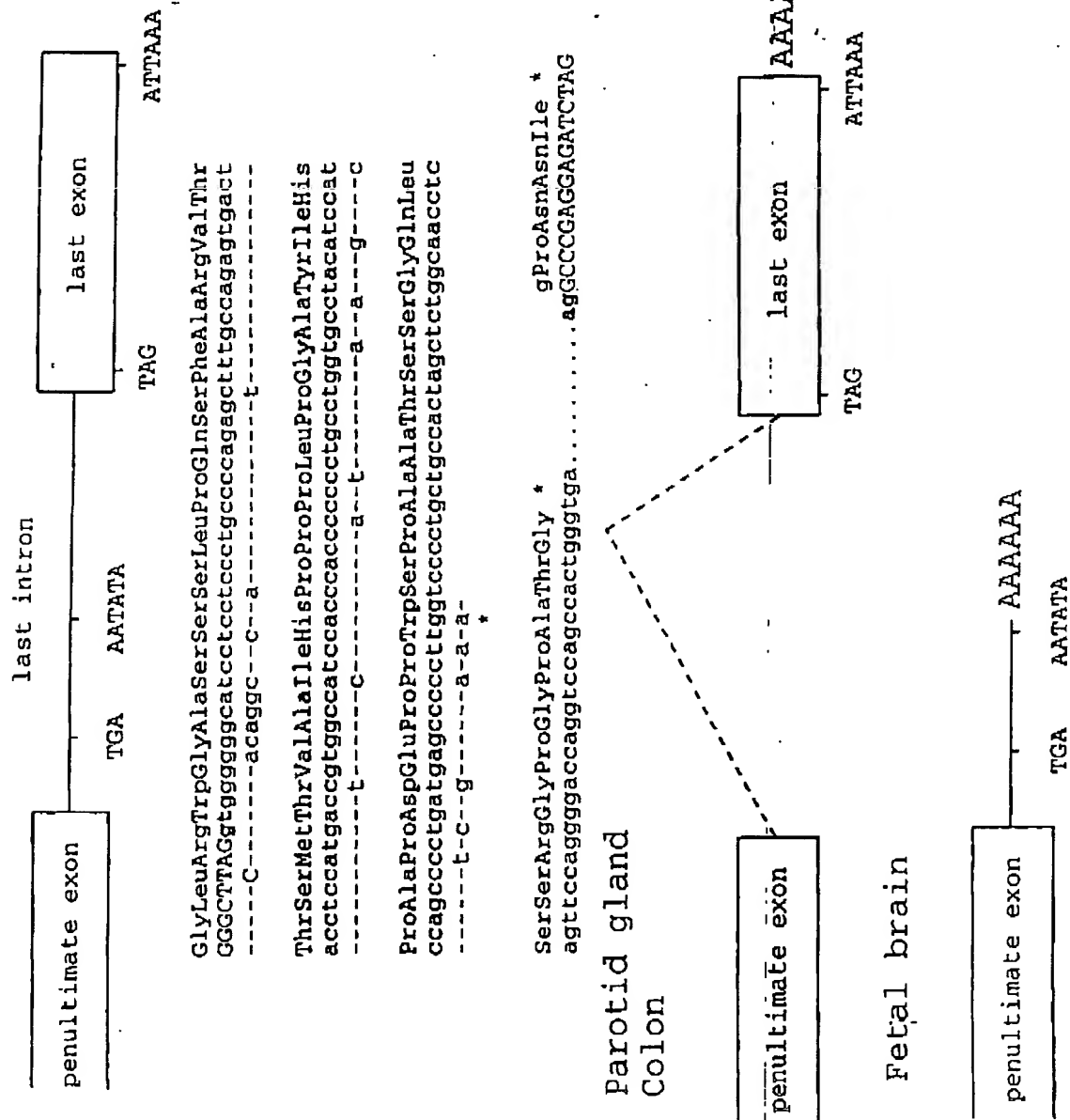
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 1 MASAGNAAEPQDRGGGSGCIGAPGRPAGGGRRRTGGLRRAAPDRDYL 50  
 12 ..PSYTPPARTAAPQI...LAGSLKAPLWLRAYFQGLLFSLGCGIQRHCG 56  
 51 HRPSYCDAA.FALEQISKGKATGRKAPLWLRKFQRLLEFKLGCIYIQKNCG 99  
 57 KVLFLGLLAFGALALGLRMAIETNLEQLWVEVGSRVSOELHYTKKLG 106  
 100 KFLVVGLLIFGAFVGLKAANLETNVEELWVEVGGRVSRELNYTRQKIGE 149  
 107 EAAVTSQMLIQATARQEGENILTPEALGLHLQAALTASKVQVSLYKSWDL 156  
 150 EAMFNPQLMIQTTPKEEGANVLTTEALLQHLDSALQASRVHVMYNNRQWKL 199  
 157 NKICYKSGVPLIENGMIERMIEKLPFCVILTPLDCFWEAKLQGG SAYLP 206  
 200 EHLCKYSGELITETGYMDQIIEYLYPCLIIITPLDCFWEAKLQSGTAYLL 249  
 207 GRPDIQWTNLDPEQLLEELGPFA.SLEGFRELLDKAQVGQAYVGRPCLHP 255  
 250 GKPLLRWTNFDPLEFLEELKKINYQVDSWEEMLNKAEVGHGYMDRPCINP 299  
 256 DDLHCPPSAENHHSRQAPNVAHELSSGCHGF SHKFMHWQEELLLGGMARD 305  
 300 ADPDCPATAPNKNSTKPLDMALVLNGGCHGLSRKYMHWQEELIVGGTVKN 349  
 306 PQGELLRAEALQSTFLLMSPROLYEHFRGDYQTHDIGWSEEQASTVLQAW 355  
 350 STGKLVSAAHALQTMFQMLTPKQMYEHFKGYEYVSHINWNEDKAAAILEAW 399  
 356 QRRFVQLAQEALPENASQQIHAFSSTTLDDILHAFSEVSAARVVGGYLLM 405  
 400 QRTYVEVVHQSVAQNSTQKVLSTTTTLDDILKSPSDVSVIKVASGYLLM 449  
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 450 LAYACLTMLRWDCSKSQGAVGLAGVLLVALSVAAGLGLCSLIGISFNAAT 499  
 456 TQVLPFLALGIGVDDVFLLAHAFTEALPG..TPLQERMGECLQRTGTSVV 503  
 500 TQVLPFLALGVGVDDVFLLAHAFSETGQNKRIPFEDRTGECLKRTGASVA 549  
 504 LTSINNMAAFLMAALVPIPALRAFSLQAAIVVGCTFVAVMLVFPAILSLD 553  
 550 LTSISNVTAFMAALIPIPALRAFSLQAAVVVVFNFAMVLLIFPAILSMD 599  
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Fig. 2A

T06020" 20020860

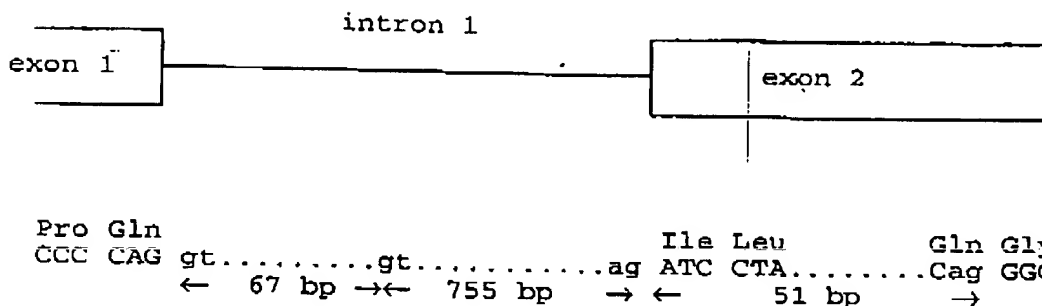
**Fig. 2B Genomic**



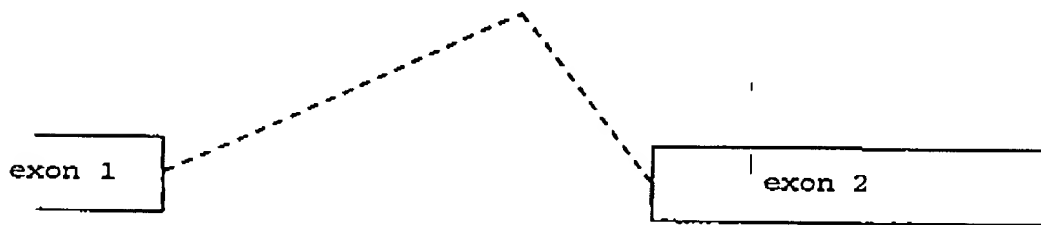
12/13

Fig. 2C

G



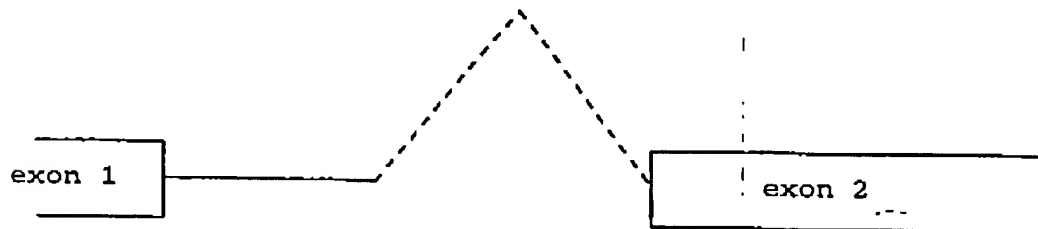
C



A



B



09807007.070901

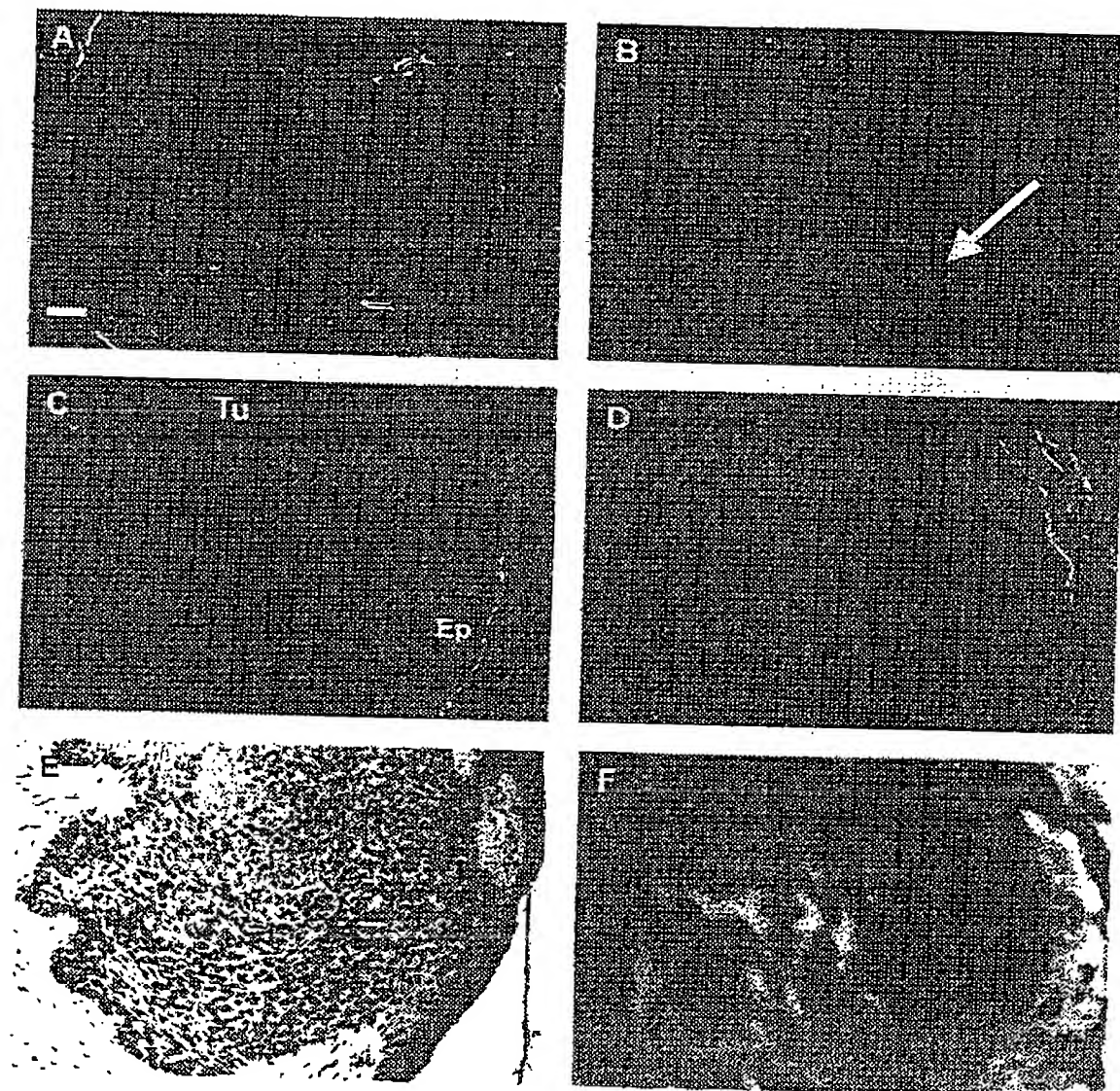


Fig. 3